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AMENDMENTS TO THE CLAIMS

1-39. (Canceled).

- 40. (Currently Amended) An isolated nucleic acid <u>encoding a polypeptide</u> having at least 80% nucleic acid sequence identity to:
- (a) a nucleic acid sequence encoding the amino acid sequence of the polypeptide of shown in Figure 4 (SEQ ID NO:9);
- (b) a nucleic acid sequence encoding the amino acid sequence of the polypeptide of shown in Figure 4 (SEQ ID NO:9), lacking its associated signal peptide;
- (c) the amino acid sequence of the polypeptide encoded by the nucleic acid sequence of shown in Figure 3 (SEQ ID NO:8);
- (d) the amino acid sequence of the polypeptide encoded by the full-length coding sequence of the nucleic acid sequence of shown in Figure 3 (SEQ ID NO:8); or
- (e) the amino acid sequence of the polypeptide encoded by the full-length coding sequence of the cDNA deposited under ATCC accession number 203406;

wherein said isolated nucleic acid encodes a polypeptide having the ability to induce c-fos expression.

- 41. (Currently Amended) An isolated nucleic acid <u>encoding a polypeptide</u> having at least 85% nucleic acid sequence identity to:
- (a) a nucleic acid sequence encoding the amino acid sequence of the polypeptide of shown in Figure 4 (SEQ ID NO:9);
- (b) a nucleic acid sequence encoding the amino acid sequence of the polypeptide of shown in Figure 4 (SEQ ID NO:9), lacking its associated signal peptide;
- (c) the amino acid sequence of the polypeptide encoded by the nucleic acid sequence of shown in Figure 3 (SEQ ID NO:8);
- (d) the amino acid sequence of the polypeptide encoded by the full-length coding sequence of the nucleic acid sequence of shown in Figure 3 (SEQ ID NO:8); or
- (e) the amino acid sequence of the polypeptide encoded by the full-length coding sequence of the cDNA deposited under ATCC accession number 203406;

wherein said isolated nucleic acid encodes a polypeptide having the ability to induce c-fos expression.

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42. (Currently Amended) An isolated nucleic acid <u>encoding a polypeptide</u> having at least 90% nucleic acid sequence identity to:

- (a) a nucleic acid sequence encoding the amino acid sequence of the polypeptide of shown in Figure 4 (SEQ ID NO:9);
- (b) a nucleic acid sequence encoding the amino acid sequence of the polypeptide of shown in Figure 4 (SEQ ID NO:9), lacking its associated signal peptide;
- (c) the amino acid sequence of the polypeptide encoded by the nucleic acid sequence of shown in Figure 3 (SEQ ID NO:8);
- (d) the amino acid sequence of the polypeptide encoded by the full-length coding sequence of the nucleic acid sequence of shown in Figure 3 (SEQ ID NO:8); or
- (e) the amino acid sequence of the polypeptide encoded by the full-length coding sequence of the cDNA deposited under ATCC accession number 203406;

wherein said isolated nucleic acid encodes a polypeptide having the ability to induce c-fos expression

- 43. (Currently Amended) An isolated nucleic acid <u>encoding a polypeptide</u> having at least 95% nucleic acid sequence identity to:
- (a) a nucleic acid sequence encoding the amino acid sequence of the polypeptide of shown in Figure 4 (SEQ ID NO:9);
- (b) a nucleic acid sequence encoding the amino acid sequence of the polypeptide of shown in Figure 4 (SEQ ID NO:9), lacking its associated signal peptide;
- (c) the amino acid sequence of the polypeptide encoded by the nucleic acid sequence of shown in Figure 3 (SEQ ID NO:8);
- (d) the amino acid sequence of the polypeptide encoded by the full-length coding sequence of the nucleic acid sequence of shown in Figure 3 (SEQ ID NO:8); or
- (e) the amino acid sequence of the polypeptide encoded by the full-length coding sequence of the cDNA deposited under ATCC accession number 203406;

wherein said isolated nucleic acid encodes a polypeptide having the ability to induce c-fos expression.

44. (Currently Amended) An isolated nucleic acid <u>encoding a polypeptide</u> having at least 99% nucleic acid sequence identity to:

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(a) a nucleic acid sequence encoding the amino acid sequence of the polypeptide of shown in Figure 4 (SEQ ID NO:9);

- (b) a nucleic acid sequence encoding the amino acid sequence of the polypeptide of shown in Figure 4 (SEQ ID NO:9), lacking its associated signal peptide;
- (c) the amino acid sequence of the polypeptide encoded by the nucleic acid sequence of shown in Figure 3 (SEQ ID NO:8);
- (d) the amino acid sequence of the polypeptide encoded by the full-length coding sequence of the nucleic acid sequence of shown in Figure 3 (SEQ ID NO:8); or
- (e) the amino acid sequence of the polypeptide encoded by the full-length coding sequence of the cDNA deposited under ATCC accession number 203406;

wherein said isolated nucleic acid encodes a polypeptide having the ability to induce c-fos expression.

- 45. (Currently Amended) An isolated nucleic acid comprising
- (a) a nucleic acid sequence encoding the amino acid sequence of the polypeptide of shown in Figure 4 (SEQ ID NO:9);
- (b) a nucleic acid sequence encoding the amino acid sequence of the polypeptide of shown in Figure 4 (SEQ ID NO:9), lacking its associated signal peptide;
- (c) the amino acid sequence of the polypeptide encoded by the nucleic acid sequence of shown in Figure 3 (SEQ ID NO:8);
- (d) the amino acid sequence of the polypeptide encoded by the full-length coding sequence of the nucleic acid sequence of shown in Figure 3 (SEQ ID NO:8); or
- (e) the amino acid sequence of the polypeptide encoded by the full-length coding sequence of the cDNA deposited under ATCC accession number 203406.
- 46. (Currently Amended) The isolated nucleic acid of Claim 45 comprising the nucleic acid sequence encoding the polypeptide of shown in Figure 4 (SEQ ID NO:9).
- 47. (Currently Amended) The isolated nucleic acid of Claim 45 comprising a nucleic acid sequence encoding the polypeptide shown in Figure 4 of (SEQ ID NO:9) lacking its associated signal peptide.

48-49. (Canceled)

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50. (Currently Amended) The isolated nucleic acid of Claim 45 comprising the nucleic acid sequence of shown in Figure 3 (SEQ ID NO:8).

- 51. (Currently Amended) The isolated nucleic acid of Claim 45 comprising the full-length coding sequence of the nucleic acid sequence of shown in Figure 3 (SEQ ID NO:8).
- 52. (Previously Presented) The isolated nucleic acid of Claims 45 comprising the full-length coding sequence of the cDNA deposited under ATCC accession number 203406.
 - 53-55. (Canceled).
 - 56. (Previously Presented) A vector comprising the nucleic acid of Claim 40.
- 57. (Original) The vector of Claim 56, wherein said nucleic acid is operably linked to control sequences recognized by a host cell transformed with the vector.
 - 58. (Previously Presented) An isolated host cell comprising the vector of Claim 56.
- 59. (Previously Presented) The host cell of Claim 58, wherein said cell is a CHO cell, an E. coli or a yeast cell.
- 60. (New) An isolated nucleic acid molecule consisting of an at least 30 nucleotide fragment of SEQ ID NO:8 or a complement thereof, that specifically hybridizes under stringent conditions to:
 - (a) the nucleic acid sequence of SEQ ID NO:8 or a complement thereof;
- (b) the full-length coding sequence of the cDNA deposited under ATCC accession number 203406 or a complement thereof;

wherein said stringent conditions use 50% formamide, 5 x SSC (0.75 M NaCl, 0.075 M sodium citrate), 50 mM sodium phosphate (pH 6.8), 0.1% sodium pyrophosphate, 5 x Denhardt's solution, sonicated salmon sperm DNA (50 μ g/ml), 0.1% SDS, and 10% dextran sulfate at 42°C, with washes at 42°C in 0.2 x SSC (sodium chloride/sodium citrate) and 50% formamide at 55°C, followed by a high-stringency wash consisting of 0.1 x SSC containing EDTA at 55°C;

and wherein said isolated nucleic acid molecule is suitable for use as a primer or a probe.

- 61. (New) The isolated nucleic acid of Claim 60 which is at least about 50 nucleotides in length.
- 62. (New) The isolated nucleic acid of Claim 60 which is at least about 75 nucleotides in length.

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63. (New) The isolated nucleic acid of Claim 60 which is at least about 100 nucleotides in length.

- 64. (New) The isolated nucleic acid of Claim 60 which is at least about 150 nucleotides in length.
- 65. (New) The isolated nucleic acid of Claim 60 which is at least about 200 nucleotides in length.
- 66. (New) The isolated nucleic acid of Claim 60 which is at least about 250 nucleotides in length.
- 67. (New) An isolated nucleic acid having at least 95% nucleic acid sequence identity to:
 - (a) the nucleic acid sequence of SEQ ID NO:8;
 - (b) the full-length coding sequence of the nucleic acid sequence of SEQ ID NO:8; or
 - (c) the full-length coding sequence of the cDNA deposited under ATCC accession number 203406;

wherein said isolated nucleic acid hybridizes to the complement of a nucleic acid of SEQ ID NO: 8 under conditions of 50% formamide, 5 x SSC (0.75 M NaCl, 0.075 M sodium citrate), 50 mM sodium phosphate (pH 6.8), 0.1% sodium pyrophosphate, 5 x Denhardt's solution, sonicated salmon sperm DNA (50 μg/ml), 0.1% SDS, and 10% dextran sulfate at 42°C, with washes at 42°C in 0.2 x SSC (sodium chloride/sodium citrate) and 50% formamide at 55°C, followed by a high-stringency wash consisting of 0.1 x SSC containing EDTA at 55°C.

- 68. (New) The isolated nucleic acid of Claim 67 having at least 99% nucleic acid sequence identity to:
 - (a) the nucleic acid sequence of SEQ ID NO:8;
 - (b) the full-length coding sequence of the nucleic acid sequence of SEQ ID NO:8; or
 - (c) the full-length coding sequence of the cDNA deposited under ATCC accession number 203406;

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wherein said isolated nucleic acid hybridizes to the complement of a nucleic acid of SEQ ID NO:8 under conditions of 50% formamide, 5 x SSC (0.75 M NaCl, 0.075 M sodium citrate), 50 mM sodium phosphate (pH 6.8), 0.1% sodium pyrophosphate, 5 x Denhardt's solution, sonicated salmon sperm DNA (50 μg/ml), 0.1% SDS, and 10% dextran sulfate at 42°C, with washes at 42°C in 0.2 x SSC (sodium chloride/sodium citrate) and 50% formamide at 55°C, followed by a high-stringency wash consisting of 0.1 x SSC containing EDTA at 55°C.

- 69. (New) A vector comprising the nucleic acid of Claim 67.
- 70. (New) The vector of Claim 69, wherein said nucleic acid is operably linked to control sequences recognized by a host cell transformed with the vector.
 - 71. (New) An isolated host cell comprising the vector of Claim 69.
- 72. (New) The host cell of Claim 71, wherein said cell is a CHO cell, an E. coli or a yeast cell.